



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/919,162 C
Source: OLPE
Date Processed by STIC: 5-7-03

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 09/914,162C

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 ☒ **Wrapped Nucleics
Wrapped Aminos** The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 ☐ **Invalid Line Length** The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 ☒ **Misaligned Amino
Numbering** The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 ☐ **Non-ASCII** The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 ☐ **Variable Length** Sequence(s) _____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 ☐ **PatentIn 2.0
"bug"** A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 ☐ **Skipped Sequences
(OLD RULES)** Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 ☐ **Skipped Sequences
(NEW RULES)** Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000
- 9 ☐ **Use of n's or Xaa's
(NEW RULES)** Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa; and which residue n or Xaa represents.
- 10 ☐ **Invalid <213>
Response** Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 ☐ **Use of <220>** Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses.
Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 ☐ **PatentIn 2.0
"bug"** Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 ☐ **Misuse of n** n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.

Does Not Comply
Corrected Diskette Needed



OIIPE

RAW SEQUENCE LISTING

DATE: 05/07/2003

PATENT APPLICATION: US/09/919,162C

TIME: 11:09:18

Input Set : N:\vernette\09919162C.RAW.txt

Output Set: N:\CRF4\05072003\I919162C.raw

1 <110> APPLICANT: Renault, Jean-Christophe
 2 Dumoutier, Laure
 4 <120> TITLE OF INVENTION: Isolated Nucleic Acid Molecules Which Encode A Soluble IL-
 TIF/IL-22
 5 Receptor or Binding
 6 Protein Which Binds to IL-TIF/IL-22, And Uses Thereof
 8 <130> FILE REFERENCE: LUD 5684.2
 10 <140> CURRENT APPLICATION NUMBER: US 09/919,162C
 C--> 11 <141> CURRENT FILING DATE: 2001-07-31
 16 <150> PRIOR APPLICATION NUMBER: US 60/245,495
 17 <151> PRIOR FILING DATE: 2000-03-11
 19 <150> PRIOR APPLICATION NUMBER: US60/234,583
 W--> 20 <151> PRIOR FILING DATE: 2000-22-09
 22 <160> NUMBER OF SEQ ID NOS: 11

ERRORED SEQUENCES

57 <210> SEQ ID NO: 5
 58 <211> LENGTH: 2271
 59 <212> TYPE: DNA
 60 <213> ORGANISM: Homo sapiens
 62 <400> SEQUENCE: 5
 63 ctgcccttaaa cccgggagtg attgtctggt tgtggatttt acagtttctt ctttggctct 60
 E--> 64 gagctgggta aaaggaacac tggttgcctg aacagtcaca cttgcaacca tgatgcctaa
 65 120 wrapped line. see item #1 on ERROR Summary SHEET
 E--> 66 acattgcttt ctaggcttcc tcatcagttt cttccttact ggtgtagcag gaactcagtc
 67 180
 E--> 68 aacgcatgag tctctgaagc ctcagagggt acaatttcag tcccgaaatt ttcacaacat
 69 240
 E--> 70 ttgcaatgg cagcctggga gggcacttac tggcaacagc agtgtctatt ttgtgcagta
 71 300
 E--> 72 caaataatat ggacagagac aatggaaaaa taaagaagac tggtggggta ctcaagaact
 73 360
 E--> 74 ctctgtgac cttaccagtg aaacctcaga catacaggaa ccttattacg ggaggggtgag
 75 420
 76 ggcggcctcg gctgggagct actcagaatg gagcatgacg ccgcggttca ctccctggtg 480
 E--> 77 ggaacaacaaa atagatcctc cagtcatgaa tataacccaa gtcaatgggt ctttgttggt
 78 540
 E--> 79 aattctccat gctccaaatt taccatatag ataccaaaag gaaaaaatg tatctataga
 80 600
 E--> 81 agattactat gaactactat accgagtttt tataattaac aattcactag aaaaggagca
 82 660
 E--> 83 aaagggttat gaaggggctc acagagcggg tgaaattgaa gctctaacac cacactccag

RAW SEQUENCE LISTING

DATE: 05/07/2003

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TIME: 11:09:18

Input Set : N:\vernette\09919162C.RAW.txt

Output Set: N:\CRF4\05072003\I919162C.raw

84 720 — Same as page 1

E--> 85 ctactgtgta gtggctgaaa tatatcagcc catgttagac agaagaagtc agagaagtga

86 780

E--> 87 agagagatgt gtggaaattc catgacttgt ggaatttggc attcagcaat gtggaaattc

88 840

E--> 89 taaagctccc tgagaacagg atgactcgtg tttgaaggat cttattttaa attgtttttg

90 900

E--> 91 tattttctta aagcaatatt cactgttaca ccttggggac ttctttgttt atccattctt

92 960

E--> 93 ttatccttta tatttcattt gtaaaactata tttgaacgac attccccccg aaaaattgaa

94 1020

E--> 95 atgtaaagat gaggcagaga ataaagtgtt ctatgaaatt cagaacttta tttctgaatg

96 1080

E--> 97 taacatccct aataacaacc ttcatctctc taatacagca aaataaaaaat ttaacaacca

98 1140

E--> 99 aggaatagta ttaagaaaa tgttgaaata atttttttaa aatagcatta cagactgagg

100 1200

E--> 101 cgttcttgaa gcaatggttt ttactctctt tattgagcca attaaattga cattgctttg

102 1260

E--> 103 acaattttaa acttctataa aggtgaatat tttcatata tttctatttt atatgaatat

104 1320

E--> 105 actttttata tatttattat tattaatat ttctacttaa tgaatcaaaa tttgttttta

106 1380

E--> 107 aagtctactt tatgtaaata agaacagggt ttggggaaaa aatcttatg atttctggat

108 1440

E--> 109 tgatatctga attaaaacta tcaacaacaa ggaagtctgc tctgtacaat tgtccctcat

110 1500

E--> 111 ttaaaagata tattaagctt ttctttctg tttgtttttg tttgttttag tttttaatcc

112 1560

E--> 113 tgtcttagaa gaacttatct ttattotcaa aattaaatgt aattttttta gtgacaaaga

114 1620

E--> 115 agaaaggaaa cctcattact caatccttct ggccaagagt gtcttgcttg tggcgcttc

116 1680

E--> 117 ctcactctta tataggagga tcccatgaat gatggtttat tgggaactgc tggggctgcac

118 1740

E--> 119 cccatacaga gaactcagct tgaagctgga agcacacagt gggtagcagg agaaggaccg

120 1800

E--> 121 gtgttggtag gtgcctacag agactataga gctagacaaa gccctccaaa ctggcccttc

122 1860

E--> 123 ctgctcactg cctctctga gtagaaatct ggtgacctaa ggctcagtgt ggtcaacaga

124 1920

E--> 125 aagctgcctt cttcacttga ggctaagtct tcatatatgt ttaaggttgt ctttctagt

126 1980

E--> 127 agagataca tatcagagaa catttgtaca attcccatg aaaattgtc caaagttgat

128 2040

E--> 129 aacaatatag tcggtgcttc tagttatatg caagtactca gtgataaatg gattaaaaa

130 2100

131 tattcagaaa tgtattgggg ggtggaggag aataagaggc agagcaagag ctagagaatt 2160

E--> 132 ggtttcttg cttccctgta tgctcagaaa acattgattt gagcatagac gcagagactg

RAW SEQUENCE LISTING

DATE: 05/07/2003

PATENT APPLICATION: US/09/919,162C

TIME: 11:09:18

Input Set : N:\vernette\09919162C.RAW.txt

Output Set: N:\CRF4\05072003\I919162C.raw

133 2220 — Same as page 1

E--> 134 aaaaaaaaaat ttactttgat ctotgttttt gaattcttat tatttatattt

135 2271

137 <210> SEQ ID NO: 6

138 <211> LENGTH: 231

139 <212> TYPE: PRT

140 <213> ORGANISM: Homo sapiens

142 <400> SEQUENCE: 6

143 Met Met Pro Lys His Cys Phe Leu Gly Phe Leu Ile Ser Phe Phe Leu

E--> 144

E--> 145 15 — misaligned Amino Acid Numbering. See item # 3 on Error Summary sheet.

146 Thr Gly Val Ala Gly Thr Gln Ser Thr His Glu Ser Leu Lys Pro Gln

E--> 147 20 25

E--> 148 30

149 Arg Val Gln Phe Gln Ser Arg Asn Phe His Asn Ile Leu Gln Trp Gln

E--> 150 35 40

E--> 151 45

152 Pro Gly Arg Ala Leu Thr Gly Asn Ser Ser Val Tyr Phe Val Gln Tyr

E--> 153 50 55 60

154 Lys Ile Tyr Gly Gln Arg Gln Trp Lys Asn Lys Glu Asp Cys Trp Gly

E--> 155 65 70 75

E--> 156 80

157 Thr Gln Glu Leu Ser Cys Asp Leu Thr Ser Glu Thr Ser Asp Ile Gln

E--> 158 85 90

E--> 159 95

160 Glu Pro Tyr Tyr Gly Arg Val Arg Ala Ala Ser Ala Gly Ser Tyr Ser

E--> 161 100 105

E--> 162 110

163 Glu Trp Ser Met Thr Pro Arg Phe Thr Pro Trp Trp Glu Thr Lys Ile

E--> 164 115 120 125

165 Asp Pro Pro Val Met Asn Ile Thr Gln Val Asn Gly Ser Leu Leu Val

E--> 166 130 135 140

167 Ile Leu His Ala Pro Asn Leu Pro Tyr Arg Tyr Gln Lys Glu Lys Asn

E--> 168 145 150 155

E--> 169 160

170 Val Ser Ile Glu Asp Tyr Tyr Glu Leu Leu Tyr Arg Val Phe Ile Ile

E--> 171 165 170

E--> 172 175

173 Asn Asn Ser Leu Glu Lys Glu Gln Lys Val Tyr Glu Gly Ala His Arg

E--> 174 180 185

E--> 175 190

176 Ala Val Glu Ile Glu Ala Leu Thr Pro His Ser Ser Tyr Cys Val Val

E--> 177 195 200 205

178 Ala Glu Ile Tyr Gln Pro Met Leu Asp Arg Arg Ser Gln Arg Ser Glu

E--> 179 210 215 220

180 Glu Arg Cys Val Glu Ile Pro

E--> 181 225 230

210 <210> SEQ ID NO: 10

211 <211> LENGTH: 2366

RAW SEQUENCE LISTING

DATE: 05/07/2003

PATENT APPLICATION: US/09/919,162C

TIME: 11:09:18

Input Set : N:\vernette\09919162C.RAW.txt

Output Set: N:\CRF4\05072003\I919162C.raw

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212 <212> TYPE: DNA
213 <213> ORGANISM: Homo sapiens
215 <400> SEQUENCE: 10
216 ctgccttaaa cccgggagtg attgtctgtt tgtggatttt acagtttcct ctttggtcct 60
E--> 217 gagctgggta aaaggaacac tggttgcctg aacagtcaca cttgcaacca tgatgcctaa
218 120 Same as page 1
E--> 219 acattgcttt ctaggcttcc tcatcagttt cttccttact ggtgtagcag gaactcagtc
220 180
E--> 221 aacgcatgag tctctgaagc ctcagagggt acaatttcag tcccgaatt ttcacaacat
222 240
E--> 223 tttgcaatgg cagcctggga gggcacttac tggcaacagc agtgtctatt ttgtgcagta
224 300
E--> 225 caaaatcatg ttctcatgca gcatgaaaag ctctcaccag agccaagtgg atgcttggca
226 360
E--> 227 gcacatttct tgtaacttcc caggctgcag aacattggct aaatatggac agagacaatg
228 420
E--> 229 gaaaaataaa gaagactggt ggggtactca agaactctct tgtgacctta ccagtgaaac
230 480
231 ctcagacata caggaacctt attacgggag ggtgagggcg gcctcggctg ggagctactc 540
E--> 232 agaattggagc atgacgccgc ggttcaactcc ctgggtgggaa acaaaaaatag atcctccagt
233 600
E--> 234 catgaatata acccaagtca atggtctctt gttggttaatt ctccatgctc caaatttacc
235 660
E--> 236 atatagatac caaaaggaaa aaaatgtatc tatagaagat tactatgaac tactataccg
237 720
E--> 238 agtttttata attaacaatt cactagaaaa ggagcaaaaag gtttatgaag gggctcacag
239 780
E--> 240 agcgggttgaa attgaagctc taacaccaca ctccagctac tgtgtagtgg ctgaaatata
241 840
E--> 242 tcagcccattg ttagacagaa gaagtcagag aagtgaagag agatgtgtgg aaattccatg
243 900
E--> 244 acttgtggaa tttggcattc agcaatgtgg aaattctaaa gctccctgag aacaggatga
245 960
E--> 246 ctctgtgttg aaggatctta tttaaaattg tttttgtatt ttcttaaagc aatattcact
247 1020
E--> 248 gttacacctt ggggacttct ttgtttatcc attcttttat cttttatatt tcatttgtaa
249 1080
E--> 250 actatatttg aacgacattc ccccccgaata attgaaatgt aaagatgagg cagagaataa
251 1140
E--> 252 agtgttttat gaaattcaga actttatttc tgaatgtaac atccctaata acaaccttca
253 1200
E--> 254 ttcttctaat acagcaaaat aaaaatttaa caaccaagga atagtattta agaaaatggt
255 1260
E--> 256 gaaataattt ttttaaaata gcattacaga ctgaggcggg cctgaagcaa tggtttttca
257 1320
E--> 258 ctctcttatt gagccaatta aattgacatt gctttgacaa tttaaaactt ctataaagg
259 1380
E--> 260 gaatattttt catacatttc tattttatat gaatataact tttatatatt tattattatt
261 1440

```

RAW SEQUENCE LISTING

DATE: 05/07/2003

PATENT APPLICATION: US/09/919,162C

TIME: 11:09:18

Input Set : N:\vernette\09919162C.RAW.txt

Output Set: N:\CRF4\05072003\I919162C.raw

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E--> 262 aaatatttct acttaatgaa tcaaaatttt gtttttaaagt ctactttatg taaataagaa
263 1500
E--> 264 caggttttgg ggaaaaaaat cttatgattt ctggattgat atctgaatta aaactatcaa
265 1560
E--> 266 caacaaggaa gtctgctctg tacaattgtc cctcatttaa aagatatatt aagctttttct
267 1620
E--> 268 tttctgtttg tttttgtttt gtttagtttt taatcctgtc ttagaagaac ttatctttat
269 1680
E--> 270 tctcaaaatt aaatgtaatt ttttttagtga caaagaagaa aggaaacctc attactcaat
271 1740
E--> 272 ccttctggcc aagagtgtct tgcttggtgc gccttctca tctctatata ggaggatccc
273 1800
E--> 274 atgaatgatg gtttattggg aactgctggg gtcgacccca tacagagaac tcagcttgaa
275 1860
276 gctggaagca cacagtgggt agcaggagaa ggaccggtgt tggtaggtgc ctacagagac 1920
E--> 277 tatagagcta gacaaagccc tccaaactgg cccctcctgc tcaactgcctc tctgagtag
278 1980
E--> 279 aaatctgggt acctaaggct cagtgtgggc aacagaaaagc tgccttcttc acttgaggct
280 2040
E--> 281 aagtcctcat atatgtttaa ggttgtcttt ctagtgagga gatacatatc agagaacatt
282 2100
E--> 283 tgtacaattc cccatgaaaa ttgctccaaa gttgataaca atatagtcgg tgcttctagt
284 2160
E--> 285 tatatgcaag tactcagtga taaatggatt aaaaaatatt cagaaatgta ttgggggggtg
286 2220
E--> 287 gaggagaata agaggcagag caagagctag agaattgggtt tccttgcttc cctgtatgct
288 2280
E--> 289 cagaaaacat tgatttgagc atagacgcag agactgaaaa aaaaatttac tttgatctct
290 2340
E--> 291 gtttttgaat tcttattatt tatattt 2366
293 <210> SEQ ID NO: 11
294 <212> TYPE: 263
W--> 295 <212> TYPE: PRT
296 <213> ORGANISM: Homo sapiens
E--> 298 <211> LENGTH:
E--> 298 <400> SEQUENCE: 11
300 Met Met Pro Lys His Cys Phe Leu Gly Phe Leu Ile Ser Phe Phe Leu
301 5 10
302 15
303 Thr Gly Val Ala Gly Thr Gln Ser Thr His Glu Ser Leu Lys Pro Gln
304 20 25
305 30
306 Arg Val Gln Phe Gln Ser Arg Asn Phe His Asn Ile Leu Gln Trp Gln
307 35 40
308 45
309 Pro Gly Arg Ala Leu Thr Gly Asn Ser Ser Val Tyr Phe Val Gln Tyr
310 50 55
311 Lys Ile Met Phe Ser Cys Ser Met Lys Ser Ser His Gln Ser Gln Val
312 65 70 75

```

Same as page 1

re-number <211>

RAW SEQUENCE LISTING

DATE: 05/07/2003

PATENT APPLICATION: US/09/919,162C

TIME: 11:09:18

Input Set : N:\vernette\09919162C.RAW.txt

Output Set: N:\CRF4\05072003\I919162C.raw

```

313 80
314 Asp Ala Trp Gln His Ile Ser Cys Asn Phe Pro Gly Cys Arg Thr Leu
315                                     85                                     90
316 95
317 Ala Lys Tyr Gly Gln Arg Gln Trp Lys Asn Lys Glu Asp Cys Trp Gly
318                                     100                                     105
319 110
320 Thr Gln Glu Leu Ser Cys Asp Leu Thr Ser Glu Thr Ser Asp Ile Gln
321                                     115                                     120                                     125
322 Glu Pro Tyr Tyr Gly Arg Val Arg Ala Ala Ser Ala Gly Ser Tyr Ser
323                                     130                                     135                                     140
324 Glu Trp Ser Met Thr Pro Arg Phe Thr Pro Trp Trp Glu Thr Lys Ile
325 145                                     150                                     155
326 160
327 Asp Pro Pro Val Met Asn Ile Thr Gln Val Asn Gly Ser Leu Leu Val
328                                     165                                     170
329 175
330 Ile Leu His Ala Pro Asn Leu Pro Tyr Arg Tyr Gln Lys Glu Lys Asn
331                                     180                                     185
332 190
333 Val Ser Ile Glu Asp Tyr Tyr Glu Leu Leu Tyr Arg Val Phe Ile Ile Asn
334                                     195                                     200                                     205
335 Asn Ser Leu Glu Lys Glu Gln Lys Val Tyr Glu Gly Ala His Arg Ala Val
336 210                                     215                                     220
337 225
338 Glu Ile Glu Ala Leu Thr Pro His Ser Ser Tyr Cys Val Val Ala Glu
339                                     230                                     235
340 240
341 Ile Tyr Gln Pro Met Leu Asp Arg Arg Ser Gln Arg Ser Glu Glu Arg
342                                     245                                     250                                     255
343 Cys Val Glu Ile Pro
344                                     260
345 IUD 5684.2 Sequences.doc -6-

```

Remove extra material at end of file.

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 05/07/2003
PATENT APPLICATION: US/09/919,162C TIME: 11:09:19

Input Set : N:\vernette\09919162C.RAW.txt
Output Set: N:\CRF4\05072003\I919162C.raw

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:6; Line(s) 144,147,150,153,155,158,161,164,166,168,171,174,179

Seq#:11; Line(s) 301,304,312,315,318,321,323,325,328,331,334,336,339,342

VERIFICATION SUMMARY

DATE: 05/07/2003

PATENT APPLICATION: US/09/919,162C

TIME: 11:09:19

Input Set : N:\vernette\09919162C.RAW.txt

Output Set: N:\CRF4\05072003\I919162C.raw

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:20 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:64 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:120 SEQ:5
M:254 Repeated in SeqNo=5
L:144 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6
M:332 Repeated in SeqNo=6
L:217 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:120 SEQ:10
M:254 Repeated in SeqNo=10
L:291 M:252 E: No. of Seq. differs, <211> LENGTH:Input:2366 Found:2367 SEQ:10
L:295 M:280 W: Numeric Identifier already exists, Type not replaced.
L:298 M:282 E: Numeric Field Identifier Missing, <211> is required.
L:298 M:310 E: (3) Wrong or Missing Sequence Type, TYPE: